

1/11

```

1  GCCACCGACA TCCGCCGCAA TGCTGTGTCT CACCTCCTCT TCCTCCTCCG CGCCCGCTCC
61  GCTCCTTCCC TCTCTCGCTG ATCGACCGAG CCCGGGAATC GCGGGCGGGG GTGGCAATGT
121  TCGCCTGAGC GTGGTTTCTT CGCCGCGCCG GTCGTGGCCT GGAAAGGTCA AGACCAATTT
181  CTCAGTTCCT GCGACTGCGC GAAAAACAA AACCATGGTG ACTGTTGTGG AGGAGGTCTGA
241  CCACCTTCCT ATATATGATC TGGACCCTAA GTTGGAGGAA TTCAAGGATC ACTTCAACTA
301  TAGGATAAAA AGATACCTCG ACCAGAAATG CCTGATTGAA AAACATGAGG GGGGCCTTGA
361  AGAATTTTCT AAAGGCTATT TGAAGTTTGG GATTAATACA GTTGATGGTG CCACAATATA
421  TCGTGAATGG GCGCCTGCTG CACAAGAAGC ACAGCTCATT GGTGAGTTCA ATAACGGAA
481  TGGTGCAAAA CACAAGATGG AGAAGGATAA ATTTGGCATT TGGTCAATCA AGATTTTACA
541  TGTCATGGG AAGCCTGCCA TCCCTCACAA TTCCAAGGTT AAATTTTCGT TTAGGCATGG
601  GGGTGGAGCA TGGGTTGATC GTATTCGCCG ATGGATTCTT TATGCAACTT TTGATGCCTC
661  TAAATTTGGA GCTCCATATG ATGGTGATCA CTGGGATCCT CCAGCCTGTG AAAGGTACGT
721  GTTTAAGCAT CCTCGACCTC CAAAACCTGA TGCTCCACGC ATCTATGAGG CTGATGTGGG
781  GATGAGTGGT GAAGAGCCAG AAGTAAGCAC ATACAGAGAA TTTGCAGACA ATGTGTTACC
841  ACGCATACGG GCAAATAACT ACAACACAGT TCAGTTAATG GCAATCATGG AACATTCCTA
901  CTATGCTTCT TTTGGGTATC ACGTGACAAA TTTTTCGCA GTCAGCAGCA GATCAGGAAC
961  ACCAGAGGAT CTGAAATATC TTGTTGACAA GGCACATAGT TTAGGATTAC GAGTTCTGAT
1021  GGATGTTGTC CATAGCCATG CGAGTAATAA TGTGACCGAT GGTCTAAATG GCTATGACGT
1081  TGGACAAAAC ACTCATGAGT CTTATTTTCA TACAGGAGAT AGGGGCTACC ATAAACTCTG
1141  GGATAGTCGT CTGTTCAACT ATGCCAATTG GGAGGTCCTA AGATTTCTTC TTTCTAATTT
1201  GAGATATTGG ATGGACGAAT TCATGTTTGA TGGCTTCCGA TTTGATGGGG TTACATCAAT
1261  GCTATACCAT CACCATGGTA TCAATAAGGG ATTTACTGGA AACTACAAGG AGTATTTTCA
1321  TTTGGATACC GATGTGGATG CAATTGTTTA CATGATGCTC GCAAACCATT TAATGCATAA
1381  ACTCTTGCCG GAAGCAACTA TTGTTGCTGA AGATGTTTCG GGCATGCCAG TGCTTTGTCTG
1441  GCCAGTTGAT GAAGGTGGAG TAGGGTTTGA CTTCCGCCTG GCAATGGCCA TTCCTGATAG
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1801  GTTTGGCCAT CCAGAAATGA TTGACTTTCC AAGAGAAGGC AACAACCTGA GCTATGATAA
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1921  ATTTGATCAA GCAATGAATG CACTCGAGGA GGAATTTTCC TTCCTGTCTAT CATCAAAGCA
1981  GATTGTTAGC GACATGAACG AGAAAGATAA GGTTATTGTC TTTGAACGTG GAGATTTGGT
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2161  AGTTGGCCAT GATGTGGATC ACTTCACGTC TCCCGAGGGA ATGCCAGGAG TACCAGAAAC
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2461  TTCTGACGAA GACTGCAATG GAAGCATCAG ATTTCTTGAT CAGGAGCAAC TGTGTTGGTC
2521  CTTGTAATCT GGAGATCCTG GCTTGCCCTG GACTTGGTTG TGGTTCTTTA GCAGTTGCTA
2581  TGTACCTATC TATGATATGA ACTTTATGTA TAGTTCGCCT TAAAGAAAGA ATAAGCAGTG
2641  ATGATGTGGC CTTAAACCTG AGCTGCACAA GCCTAATGTA AAAATAAAGT TTCAGGCTTT
2701  CATCCAGAAT AAAACAGCTG TTCATTTACC ATCTCAAAA

```

Figure 1

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1 CTTGACTCCC CCCACTCCTC CCTCGTGCTG CTCCTCCTCG TCGCTCGGCT CGAGGCGCGG
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181 GCGGCGGCGG CGGCGGGGGT GGCCCGGCGG CGCGATCCGG CGGGGTGGAC TTGCCGTCGG
241 TGCTCTTCAG GAGGAAGGAC TCCTTCTCAC GTGGCGTTGT GAGCTGCGCG GGTGCTCCTG
301 GGAAGGTGCT GGTGCCGTGC GGTGGGAGCG ACGACTTGCT GTCTCTGCG GAACCAGACG
361 TGGAACTCA AGAGCAACCT GAAGAATCTC AGATACCTGA TGATAATAAA GTAAAACCTT
421 TTGAGGAGGA GGAAGAGATT CCAGCAGTGG CAGAAGCAAG CATAAAGGTT GTGGCTGAAG
481 ACAAACTTGA ATCTTCAGAA GTGATTCAAG ACATTGAGGA AAATGTGACT GAGGGTGTGA
541 TCAAAGATGC TGATGAACCA ACTGTGGAGG ATAAACCACG AGTTATCCCA CCACCAGGAG
601 ATGGGCAGAA GATATACCAA ATTGACCCAA TGCTGGAAGG ATTTCCGAAC CATCTTGACT
661 ACCGATACAG TGAATACAAG AGAATGCGTG CAGCTATTGA CCAACATGAA GGTGGCTTGG
721 ATGCATTTTC TCGTGGTTAC GAAAAGCTTG GATTCAACCG CAGCGCTGAA GGCATTACCT
781 ACCGAGAATG GGCACCTGGA GCACAGTCTG CAGCATTAGT AGGTGACTTC AACAAATTGA
841 ACCCAAATGC AGATACTATG ACCAGAAATC AGTATGGTGT TTGGGAGATT TCCTGCCTA
901 ACAATGCTGA TGGATCCCTT GCTATTCCCT ATGGCTCAGG TGTAAGATT CGGATGGATA
961 CACCATCTGG CGTAAAGGAT TCAATTCCTG CCTGGATTAA GTTTGCTGTG CAGGCTCCAG
1021 GTGAAATACC GTACAACGGT ATATATTATG ATCCACCTGA AGAAGAAAAA TATGTATTCC
1081 AACATCCTCA ACCTAAACGA CCAAATTCGC TGCGGATATA TGAATCACAT ATTGGAATGA
1141 GTAGCCCGGA ACCGAAGATA AACACATATG CTAATTTTAG GGATGAGGTG CTACCAAGAA
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1261 CAAGCTTTGG GTATCATGTT ACTAACTTCT TTGCGCCAAG TAGCCGTTTC GGAACCCAG
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1501 TCAACTATGG GAGTTGGGAA GTTTTAAGAT ATTTACTGTC GAATGCAAGG TGGTGGCTTG
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1621 ATGGTTTACA GGTGGCATT ACTGGCAACT ATGGCGAATA TTTTGGATTT GCTACTGATG
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1981 AGACTATTGC ATTCTGGTTG ATGGATAAGG ATATGTATGA TTTTATGGCT CTAGACAGAC
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2101 TGGGCTTAGG AGGCGAAGGC TATCTTAATT TCATGGGAAA TGAGTTTGGG CATCCTGAAT
2161 GGATAGATTT CCAAGAGGC CCGCAAAGTC TTCCAATGG CTCGGTCTCT CCAGGAAACA
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2641 TTACAGAGGA CTAATGATCA GCTCTGATCA TTGGGGGAAC AACTCAAGGG AGTTGGTGGT
2701 AATGACGCCG GAATACAACT CAAGTGAAAG GTGAAAAGAA AGGCTGCCCT GACGATGTGA
2761 TTTGAGGGGC TTGTGTTTCA TCGCCAATGC CAGGAAGATG AGGTAGAAAA GCCTACTGAT
2821 GAGCTCCTGT TTTGAGTGA CTCGTGAAGG AAATAGACCA GGGTGAACGG CTTTTTTCAG
2881 AGCTATACCA AACCATCCTT ATGTTGCGCA TTCGCTGTAG TTTTGTACAT AACGATATCG
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Figure 2

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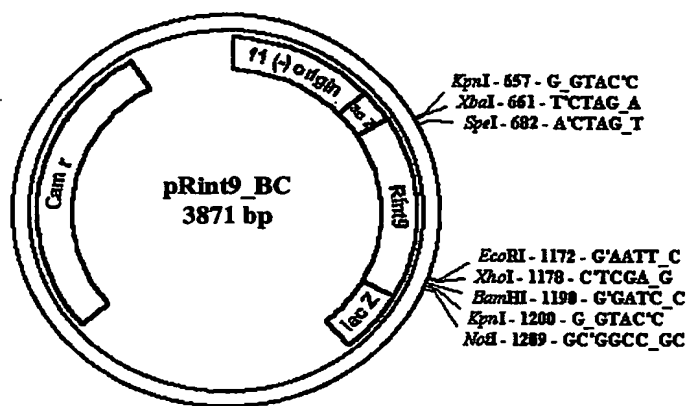
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121 GGGCGAGATG GCGGCGCCGG CGTCTGCGGT TCCCGGGAGC GCGGCGGGGC TACGGGCGGG
181 GGCCGTGCGG TTCCCGGTGC CAGCCGGGGC CCGGAGCTGG CGTGGCGGGC CGGAGCTCCC
241 GACGTGCGCG TCGCTGCTCT CCGGCCGAG ATTCCCCGGT GCCGTTTCGG TGGGGGGTTC
301 CGGGGGGCGC GTGGCCGTGC GCGGCGGGG CGCGTCAGGG GAGGTGATGA TCCCCGAGGG
361 CGAGAGCGAC GGGATGCCGG TTTCAGCAGG TTCAGACGAT CTGCAGTTGC CAGCCTTAGA
421 TGATGAATTA AGCACGGAGG TTGGAGCTGA AGTTGAGATT GAGTCATCTG GAGCAAGTGA
481 CGTTGAAGGC GTGAAGAGAG TGGTTGAAGA ATTAGCTGCT GAGCAGAAAC CACGAGTTGT
541 CCCACCAACA GGAGATGGGC AAAAAATATT CCAGATGGAC TCTATGCTTA ATGGCTATAA
601 GTACCATCTT GAATATCGAT ATAGCCTATA TAGGAGACTG CGTTCAGACA TTGATCAGTA
661 TGAAGGAGGA CTGGAACAT TTTCGCGGG TTATGAGAAG TTGGATTATA ATCAGAGTGC
721 TGAAGGTGTC ACTTATCGAG AATGGGCTCC CGGGGCGACAT TCTGCAGCAT TAGTAGGTGA
781 CTTCACAAT TGAATCCAA ATGCAGACCG CATGAGCAAA AATGAGTTTG GTGTTTGGGA
841 GATTTTTCTG CCTAACAATG CTGATGCTC ATCTCCTATT CCACATGGCT CACGTGTAAA
901 GGTGCGAATG GAACTCCAT CTGGTATAAA GGATTCATT CCTGCCTGGA TCAAGTACTC
961 TGTGCAGGCC GCAGGAGAAA TCCCATACAA TGGAATATAT TATGATCCTC CTGAAGAGGA
1021 GAAGTACATA TTCAAGCATC CTCAACCTAA AAGACCAAAG TCATTGCGGA TATACGAAAC
1081 TCATGTTGGA ATGAGTAGCA CGGAGCCAAA GATCAACACG TATGCAAACT TTAGGGATGA
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2041 TAGACTTATC ACAATGGGGT TAGGAGGAGA AGGCTATCTT AACTTTATGG GAAATGAGTT
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2161 CATCCAGGG AATAACAACA GTTATGATAA ATGCCGTCGA AGATTTGACC TGGGTGATGC
2221 GGAATATCTT AGGTATCGTG GCATGCTAGA GTTTGACCGC GCGATGCAGT CTCTCGAGGA
2281 AAAATATGGG TTCATGACAT CAGACCACCA GTACATATCT CGAAAGCATG AAGAGGATAA
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2701 TCTGCTTCGA TGAATGCCGG ATAGACTAGA CAGCTTGCTT TTGTGCTTTG CGCTCCCAAT
2761 TTGTAGTTTT AGTTTGTGAG GGAAAGAAAC GTTTATTTGT AATTATCTAT GGCTGTCGAA
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Figure 3

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**Figure 4**

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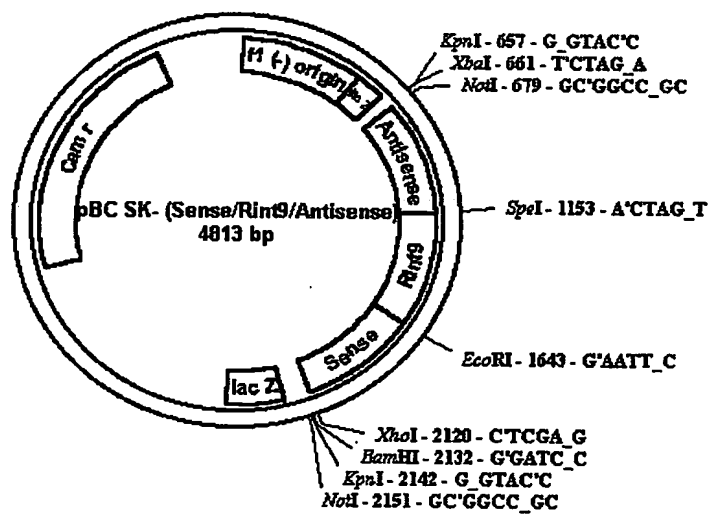
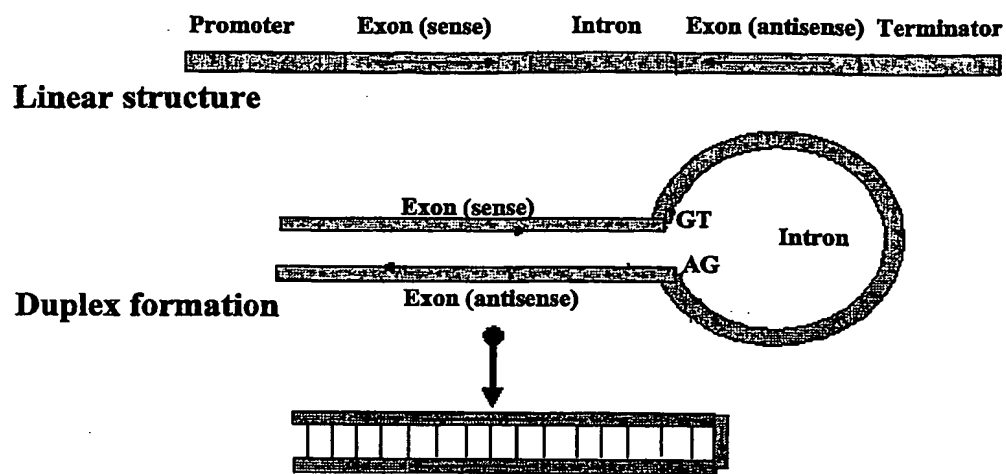


Figure 5

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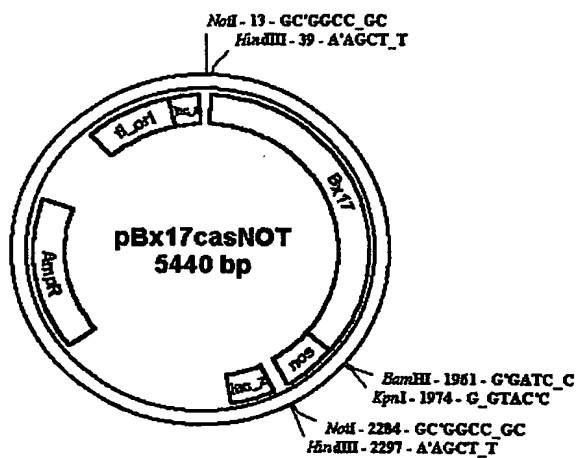


Figure 6

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55  gcgcggcatttgcggcggga.gggatctgcgcgcgagtgcggtcgggcag 103
    .
131 GCGGC.....GCCGGCGTCTGCGGTTCCTCGGGA 158
    |||||  |||||  |||||  |||||  |||||
104 gcggcgggggagcacgcaccgggggatggcgctcgttcgcggtgtcc.ggc 152
    .
159 GCGCGGGGGGGCTACGGGCGGGGGCCGTGCGGTTCCTCGCAGCCGGG 208
    ||  |||||  ||  |||||  ||  |||||  ||  |||||
153 gcgaggtcgggggtcgtgcgggcggggggcgg....cggcggcggcgggg 198
    .
209 GCCCGGAGCTGGCGTGCGGCGGGCGGAGCTCCCGACGTGCGGTTCGCTGCT 258
    |  |||||  |  |||||  |  |||||  |  |||||
199 gtggcccgggcgcgcatccggcgggg....tggacttgccgtcggtgtct 244
    .
259 CTCCGGCCGGAGATTCCCGGTGCCGTTCCGCTGGGGGGTTCGGGGGGG 308
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245 cttcaggagga.....aggactccttctcacgtggcggt..... 278
    .
309 GCGTGGCCGTGCGCGCGGGCGGCGTCAGGGGAGGTGATGATCCCGAG 358
    |||||  |||||  |||||  |||||  |||||
279 .....gtgagctgcgcgggtgctcctgggaagggtgctggtgcctggc 320
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359 GGCGAGAGCGACGGGATGCCGGTTTCAGCAGGTTTCAGACG..... 398
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321 ggtgggagcgacgacttgctgtcctctgcggaaccagacgtggaaactca 370
    .
399 .....ATCTGCAGTTGCC.....AGCCT 416
    |||||  |||||  |||||
371 agagcaacctgaagaatct.cagataacctgatgataataaagtaaacct 419
    .
417 T.....AGATGATGAATTAAGCACGGAGGT 441
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420 tttgaggaggaggaagagattccagcagtggcagaagcaagcataaagg 469
    .
442 TGGAGCTGAAGTTGAGATTGAGTCATC.....TGGAG 473
    ||  |||||  |||||  |||||
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    .
474 CAAGTGACGTTGAAGGCGTGAAGAGAGTGGTTGAAGAATTAGCTGCTGAG 523
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520 aaaatgtgactgaggggtgatcaaatgctgatgaaccaactgtggag 569
    .
524 CAGAAACCACGAGTTGTCCACCAACAGGAGATGGGCAAAAAATATTCCA 573
    |  |||||  |||||  |||||  |||||
570 gataaaccacgagttatcccaccaccaggagatgggcagaagataatacca 619
    .
574 GATGGACTCTATGCTTAATGGCTATAAGTACCATCTTGAATATCGATATA 623
    ||  |||||  |||||  |||||  |||||
620 aattgaccacaatgctggaaggatttcggaaccatcttgactaccgatata 669
    .
624 GCCTATATAGGAGACTGCGTTTCAGACATTGATCAGTATGAAGGAGGACTG 673
    |  |||||  |||||  |||||  |||||
670 gtgaataacaagagaatgcgtgcagctattgaccaacatgaagggtggcttg 719

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Figure 7

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    || ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
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    .
724 AGGTGTCACCTATCGAGAATGGGCTCCCGGGGCACATTCTGCAGCATTAG 773
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    .
774 TAGGTGACTTCAACAATTGGAATCCAAATGCAGACCGCATGAGCAAAAAT 823
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820 taggtgacttcaacaattggaacccaaatgcagatactatgaccagaaat 869
    .
824 GAGTTTGGTGTGTTGGGAGATTCTTCTGCCTAACAATGCTGATGGCTCATC 873
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870 gagtatggtggttgggagatttccctgcctaacaatgctgatggatcccc 919
    .
874 TCCTATTCACATGGCTCACGTGTAAAGGTGCGAATGGAACTCCATCTG 923
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920 tgctattcctcatggctcacgtgtaaagattcggatggatacaccatctg 969
    .
924 GTATAAAGGATTCTATTCCCTGCCTGGATCAAGTACTCTGTGCAGGCCGCA 973
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970 gcgtaaaggattcaattcctgcctggattaagtttgcgtgtgcaggctcca 1019
    .
974 GGAGAAATCCCATACAATGGAATATATTATGATCCTCCTGAAGAGGAGAA 1023
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1020 ggtgaaataccgtacaacggtatatattatgatccacctgaagaagaaaa 1069
    .
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    .
1074 ACGAAACTCATGTTGGAATGAGTAGCACGGAGCCAAAGATCAACACGTAT 1123
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    .
1124 GCAAACCTTAGGGATGAGGTGCTTCCAAGAATCAAAAAGCTTGGATACAA 1173
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1174 TGCAGTGCAAATAATGGCAATTCAAGAGCATGCATATTATGGAAGCTTTG 1223
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1224 GGTACCATGTCACCAATTCTTTGCACCAAGTAGTCGTTTCGGGACCCCA 1273
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1270 ggtatcatgttactaacttctttgcgccaagtagccgtttcggaacccca 1319
    .
1274 GAAGATTTAAAGTCATTGATTGATAAAGCTCATGAGCTTGGTTTAGTTGT 1323
    |||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
1320 gaagacttgaaatctctgattgataaagctcacgagcttggtttgcctgt 1369
    .
1324 GCTCATGGATGTTGTTTACAGCCATGCGTCAAATAATACCCTAGATGGGT 1373
    || ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
1370 acttatggatattgttcacagtcatgcatcaaacaataccctggatgggtt 1419

```

Figure 7

| | | |
|------|---|------|
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| 1420 | tgaatggttttTgatggtactgatacacattacttccatggTggaccacgg | 1469 |
| 1424 | GGCCATCATTTGGATGTGGGATTCTCGCCTTTTCAACTATGGGAATTGGGA | 1473 |
| 1470 | ggtcatcactggatgtgggattctcgctgttcaactatgggagttggga | 1519 |
| 1474 | AGTTCTAAGATTTCTACTATCCAATGCAAGATGGTGGCTCGAGGAGTATA | 1523 |
| 1520 | agttttaagatatttactgtcgaatgcaaggTggtggcttgaagaataca | 1569 |
| 1524 | AGTTTGATGGTTTCAGATTTGACGGTGTAACCTCAATGATGTACACTCAT | 1573 |
| 1570 | agtttgatgggtttcgatttgatggggTgacctccatgatgtatactcat | 1619 |
| 1574 | CATGGATTACAAGTAGCATTTACGGGGAAC TACAGTGAATACTTTGGATT | 1623 |
| 1620 | catggtttacaggTggcatttactggcaactatggcgaatattttggatt | 1669 |
| 1624 | TGCCACTGATGCTGATGCAGTAGTTTACTTGATGCTGGTAAATGATT TAA | 1673 |
| 1670 | tgctactgatgttgatgcagtagtttacttgatgctggtgaacgatctaa | 1719 |
| 1674 | TTCATGGACTTTATCCTGAGGCCATAACCATCGGTGAAGATGTCAGTGGA | 1723 |
| 1720 | ttcatgggctttatcctgaggctgtagccattggtgaagatgtcagcggg | 1769 |
| 1724 | ATGCCTACATTTGCCCTTCTGTTC AAGATGGTGGGGTTGGTTTTGATTA | 1773 |
| 1770 | atgccacattttgtattcctgttcaagatggTggtggttggttttgacta | 1819 |
| 1774 | TCGCCTTCATATGGCTGTTTCTTGACAAATGGATTGAACTCCTCAAGCAA | 1823 |
| 1820 | tcgtttgcatatggctgtaccggacaaatggatcgaaactcctcaagcaa | 1869 |
| 1824 | GTGATGAATCTTGAAGATGGGTGATATTGTGCACACACTGACTAACAGA | 1873 |
| 1870 | gtgacgaatatttgaaaatgggtgatatcgtgcacaccctaacgaataga | 1919 |
| 1874 | AGGTGGTCAGAGAAGTGTGTTACTTATGCTGAAAGTCATGATCAAGCACT | 1923 |
| 1920 | aggtggtcagagaagTgtgttacttatgcagaaagtcatgaccaagcact | 1969 |
| 1924 | AGTTGGTGACAAAAC TATTGCATTCTGGTTGATGGACAAGGATATGTATG | 1973 |
| 1970 | agttggTgacaagactattgcattctgggtgatggataaggatatgtatg | 2019 |
| 1974 | ATTTTATGGCTCTGGACAGACCGGCAACACCTAGCATTGATCGTGG AATA | 2023 |
| 2020 | attttatggctctagacagaccttcaacacctcgcatTgatcgtgggata | 2069 |
| 2024 | GCATTGCATAAAATGATTAGACTTATCACAATGGGGTTAGGAGGAGAAGG | 2073 |
| 2070 | gcattacataaaatgattaggctgtcaccatgggcttaggagggcqaagg | 2119 |

Figure 7

Figure 7

11/11

riceSBEIIaIR.seq

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1      CTCGAGTCTA  GATCGCGTC  G GTTGTTTA  AA GCCTGGA  AAG TACAAG  ATTGT
56     GTTGGACTC  AGACGATGGC  CTCTTTGGT  G GATTCACT  CG GCTTGAT  CATGA
111    TGCTGAGT A  CTTCACTGC  TGACTGGCCG  CATGACAAC  A GACCATGT  TCATT
166    CTCGGTG TA  CACCCCAA G  CAGAACCGC  CGTCGTGTAT  GCACTTACA  GAGGA
221    CTAATG ATC  AGCTCTG AT  CATTGGGG G  AACAACCTCA  AGGGAGTTGG  TGGTA
276    ATGAC GCCG  GAATAC AAC  TCAAGTG AA  AGGTGAAA A  GAAAGGCTGC  CCTGA
331    CGAT GTGAT  TTGAG GGGC  TTGTGT TTC  ATCGCCA AT  GCCAGGAAGA  TGAGG
386    TAG AAAAGC  CTAC TGATG  AGCTC CTGT  TTTCGA GTG  ACTCGTGAAG  GAAAT
441    AG ACCAGGG  TGA ACGGCT  TTTT TCAGA  GCTAT ACCA  AACCACATCCT  ATGTT
496    G CGCATTCG  CT GTAGTTT  TGT ACATAA  CGAT ATCGG  TTGGCATTG  TATGT
551    TTATGAATA  A TCTGTTCG  AC AGAAATG  TTT TTCTCC  TTGTAAC TAG  TGAA
606    TTC

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riceSBEIIbIR.seq

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1      CTCGAGTCTA  GNNNNNNNNN  N NNNNNNNN  NN NNNNNNN  NNN NNNNNN  NNNNN
56     NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  N NNNNNNNN  NN NNNNNNN  NNNNN
111    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNN  N NNNNNNNN  NNNNN
166    NNNNNNNN NN  NNNNNNNNN  N NNNNNNNNN  NNNNNNNNNG  CTCCAGCGG  AATGA
221    GAACAC CAA  GAGGCAG CA  TGCAAGTG T  GTGCGGCTG  CTAGTGC GAA  GGAGC
276    AAGAA AAAC  TAGTTG CCA  GCAATCT GT  GAACGGCT T  TCCTAGGTTT  TGCTT
331    CGAT GAATG  CCGGA TAGA  CTAGAC ANN  NNNNNNNN NN  NNNNNNNNNN  NNNNN
386    NNT TGTAGT  TTTA GTTTG  TGAGG GAAA  GAAACG TTT  ATTTGTAATT  ATCTG
441    TG GCTGTCG  AAC GGCGAC  GAAA CCATG  AACCC CGTA  TATTTGTTGG  TACCG
496    T TCGAACTG  CC AGTTATA  CAT AGTTCT  GCAC TTCTG  TACATCTTGT  GATGC
551    TACTAGTGA  A TTC

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riceSBEIIR.seq

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1      CTCGAGTCTN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
56     NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
111    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
166    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
221    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
276    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
331    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
386    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
441    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
496    NNNNNNNNNN  NNNNNNNNAGC  ATCAGATTTC  TTGATCAGGA  GCAACTGTTG  GTGCC
551    CTTGTAAACT  AGTGAATTC

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Figure 8